

#6

RAW SEQUENCE LISTING DATE: 10/20/2000
PATENT APPLICATION: US/09/557,796 TIME: 11:48:15

Input Set : A:\557796.txt
Output Set: N:\CRF3\10202000\I557796.raw

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65	cagatttca	tcgttattgt	atggataag	ctgatcgacg	gccaacccg	cgacgccaag	840
66	ctgcacgga	ttatggatt	catcaccacc	gcccggcg	ccgatgaaa	tgtggcggtc	900
67	cgtcttctg	gcatgaatt	ttcccgctc	ctggatgaaa	accggccaa	cgccattacc	960
68	gtcgatgaca	gcatgtggc	caaaattcag	gcccgt			996
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72	<211>	LENGTH:	462				
73	<212>	TYPE:	DNA				
74	<213>	ORGANISM:	yia.1				
76	<400>	SEQUENCE:	3				
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79	gcccgttatt	tcctcgccac	gacggatttc	cacgcgttgc	caccggcg	cgtggaaatc	120
80	gacggccaaa	acatcttcgc	gcagggttac	gacttaacca	ctcgcatgc	cgctgaaaat	180
81	cgtccggagg	tcacccgtcg	ctatcttgc	atccgtttc	tggcatcg	cgaagaaaaa	240
82	atcggtatcg	ccattgatac	cggcaataat	caaattcagc	aatctttatt	agaacagcgc	300
83	gatattttt	tttatcaca	cagcgaacat	gaatcg	ttgaaatgac	gccaggcaac	360
84	tatgcgat	ttttccgc	agatgttcat	ctgcctggat	gtataaaaaac	tgtagccacg	420
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91	<213>	ORGANISM:	orf1				
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96	tcatttacc	aaaagagtga	agaggaaagag	atggaaattt	ggcggcagct	ctccgacacc	120
97	ccggatattc	gccccgttgc	gcaaccctgc	ctaccctgc	ttgaacat	tcatccgtc	180
98	ggcgcacgt	gttattcgc	ccatcccccg	ggacactggc	agatgtcg	taccgcatac	240
99	atggaaacca	tgcggccgc	cgtgaaaac	ggcggcttgc	ggctggcg	cagcgcacaa	300
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101	aatggcaata	ccggccggaa	ggtcatttgc	cttgcatttgc	acggccgc	gctggccggc	420
102	aatggcaacg	tggctcaggc	tacccgc	tttgccttgc	cattaaa	aatttccgc	480
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104	cgcacaggat	tttgcgtt	agaaaacgt	cttgcgc	tttgcgat	tgacgttgc	600
105	atttgcattt	actggccgc	tttgcgc	gaaggccgc	ataccgtgt	accgcgc	660
106	catacgc	aggtaaaacg	ggcaggaaag	ctccggcg	tgatgtttc	tggcgcacg	720
107	cagacccgc	agtaacgc	atgcgcaggat	ttacacgc	cgatgcgc	tttgcgc	780
108	cagacccgc	tgaccaccgc	acacgcgt	gaattatttgc	cgtgcgc	agg aaccgc	840
109	ctgcaattt	caggcattaa	attactggaa	attatgc	gcaaaacgt	tgcgc	900
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115	<212>	TYPE:	DNA				
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121	attccgccta	tactgtatc	ttgttattt	tcttatatgg	accgggttca	tattgc	120
122	gcatgcgc	gaggatgttgc	ttccgcactt	ggtatttgc	ccaccatgc	ggggctggcg	180
123	ggcggtattt	tcttatatgg	ttatcttattt	ttacagg	ccggccggaa	aattgcgtt	240
124	cacgttagcg	gtaaagaaat	tatccgc	tcgcgttgc	cctggccgtt	cattccgt	300
125	ctgacggggt	taattacaa	tcagtc	ctgcgtggcc	tgcgtt	actggcg	360

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126	gcggaaaggcg	gtatgtgcc	ggtcgttctc	acgatgtatca	gtaaactgggtt	ccccgacgct	420
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128	gccccactct	caggctggat	tatcacgggtt	ctcgactggc	gttggctgtt	tattatcgaa	540
129	ggtttgcgtt	cgctgggtt	tctgggtctg	tgggcataca	ccatctatga	ccgtccgcag	600
130	gaaggcgcgt	ggatttccga	agcagagaag	cgctatctgg	tcgagacgt	ggccgcggag	660
131	caaaaagcca	ttggccggcac	cgaggtgaaa	aacgcctctc	tgagcgcgt	tctctccgac	720
132	aaaaccatgt	ggcagcttat	cgccctgaac	ttcttcttacc	agacccggat	ttacggctac	780
133	accctgtggc	taccacatc	tctgaaagaa	ttgaccatca	gcagcatgg	gcaggtcggc	840
134	atgtttgcga	ttctgcgtta	cgtccggcgc	attgtcggtt	tgttccctgtt	ttctccctt	900
135	tcagaccgaa	cggtaaaacg	caagctgttc	gtctgectgc	cgctgattgg	cttcgccttg	960
136	tgcattgtcc	tgtcggtggc	gtcgaaaaaac	caaatttggc	tctcctatgc	cgcgctggc	1020
137	ggctcgggat	tcttcctgc	atcgccggct	ggcattgtttct	ggaccatccc	ggcacgtctg	1080
138	ttcagcgcgg	aaatggcggg	cggcgcgcgc	ggggttatca	acgcgttgg	caaccccgcc	1140
139	ggattttgtt	gcccattatgc	gttcgggggtt	ctgatcacgt	tgtacagcaa	agacgttggc	1200
140	gtctattggc	tggcgatctc	cctggcgctg	gccgcgtga	tggcgctgt	gtcgccggcg	1260
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147	<213> ORGANISM: lyxk						
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153	tcgccccctgc	cgggttacgc	cgaacgcgcac	atgcgcacgc	tctggcaaca	ctgcgcggcg	180
154	accattgtcc	ggcttattaca	cgaggcgagg	gtatccggcg	aacagattaa	aggcggtggc	240
155	atctccgtc	agggtcaagg	gcttttctc	ctcgataacg	aggatcgcc	gttgggttaac	300
156	gccatctct	cctccgtatcg	tcggcgctg	aaaatcgttc	agcgctggca	gcgggaccgt	360
157	atccccgaac	ggctctatcc	cgttaccgc	cagacgtgt	ggaccggaca	tccggcttct	420
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162	tgcggggaga	tcaccgtca	ggcagccgt	ttaacccgtc	tggcgccggg	tactcccg	720
163	gttggcgccc	tgtttgacat	ttgttgcacc	gcccatttgc	cggggattga	ggatgagtgc	780
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166	cacaaaggca	gcccggaccc	atccggcaac	ctcgaaatgtt	ttaccgc	gtggggcgat	960
167	ctctcggtt	atgagatcaa	tcaggccgtc	gccagectgc	cgaaaggcgg	gaggcgacgt	1020
168	tttttctgc	cgtttctgt	tggcagcaac	gcccggctgg	agatgaccc	cggttttac	1080
169	ggcatgcagg	cgctgcatac	ccgcgcgcac	ctgctgcagg	cggtttatga	aggcggtgg	1140
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173	gctcggtcg	gtaccggcgt	ataccgcgc	tttagcgaag	ccggcgccgc	ccggcagcac	1380
174	ccgggtcgca	cgctgtgc	cgatatgacc	gcccacgcgc	gtatcg	caaataccgc	1440
175	cactacctgc	atttgcgttgc	agcactacag	ggctatc	cccgatattaa	ggagcacgc	1500
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180	<211> LENGTH: 660						

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182 <213> ORGANISM: yia q
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188 accgaaggc tttagcgcggg taaaggccctg cgccggccagg gtccggggaa gatcatcg
189 gccactgga aatgcgcgca cggccgtgaa acccttggcc agcaggccctt tggcgctgc
190 gccaactgga tgaccatcat ttgcgcgca cggccgtgca cggccgtgc
191 gtggcccgagg cttgcggcggg taaatttcgat atggagctgt tcggcaactg gacgttgat
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194 gatattggcc tttagcttgc gattaccggc ggcatttaccc cagccgtatcc acccgctgtt
195 aaagatataca acgtcaaaacg cttttttgc gggcgcgcgc tggcaggcgc cggccatccg
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202 <213> ORGANISM: yia r
204 <400> SEQUENCE: 8
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210 cccttggca gcccgcgttgc aeggttgcgtc gatcggggcgc gagagattat gacccaaaggc
211 atccgcgttgc cggcgcgtatcc ggggttgcaccatccacg tggcgggttgc cggatgttca
212 tagcggaaaggc atgtatggg caccggcggcgttgc aagggttgcggc aagggttgcgc
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218 aagacgttgc gcgagactgaa ctaccggcgttgc tcatgttgc gtttaccgtc
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224 <211> LENGTH: 714
225 <212> TYPE: DNA
226 <213> ORGANISM: yia s
228 <400> SEQUENCE: 9
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238 ggcacgttcc cggccgttgc gttccatca caccggccctt tggcgggggaaa aagacgc
239 gcccggccgc tacataacgc cttggatgtcg gggatgtcg cttccatgttgc cttcttcg
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540
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660
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780
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858

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240 cggcagtggc cacagctgcc ggatatgcag tctgaactgc tcgataaaca ctatctgcgt 660
241 aaacacggcg cgaacgctat tacgggcaaa actagtcccg cggaactccc cgga 714
244 <210> SEQ ID NO: 10
245 <211> LENGTH: 272
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249 <400> SEQUENCE: 10
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252 1 5 10 15
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255 20 25 30
257 Asn Tyr Pro Asn Gly Cys Pro Val Ala His Leu Ser Glu Leu Ala Gly
258 35 40 45
260 Leu Asn Lys Ser Thr Val His Arg Leu Leu Gln Gly Leu Gln Ser Cys
261 50 55 60
263 Gly Tyr Val Thr Pro Ala Ala Gly Ser Tyr Ala Leu Thr Thr
264 65 70 75 80
266 Lys Phe Ile Arg Val Gly Gln Lys Ala Leu Ser Ser Leu Asn Ile Ile
267 85 90 95
269 His Val Ala Ala Pro His Leu Glu Ala Leu Asn Leu Ala Thr Gly Glu
270 100 105 110
272 Thr Val Asn Phe Ser Ser Arg Glu Asp Asp His Ala Ile Leu Ile Tyr
273 115 120 125
275 Lys Leu Glu Pro Thr Thr Gly Met Leu Arg Thr Arg Ala Tyr Ile Gly
276 130 135 140
278 Gln His Met Arg Cys Thr Ala Arg Gln Trp Ala Lys Ile Tyr Met Ala
279 145 150 155 160
281 Phe Gly His Pro Asp Tyr Val Glu Ser Tyr Trp Asn Ser His Gln Glu
282 165 170 175
284 Ile Ile Gln Pro Leu Thr Arg Asn Thr Ile Thr Gly Leu Pro Ala Met
285 180 185 190
287 His Asp Glu Leu Ala Gln Ile Arg Glu Arg Asn Met Ala Met Asp Arg
288 195 200 205
289 Glu Glu Asn Glu Leu Gly Val Ser Cys Leu Ala Val Pro Val Phe Asp
290 210 215 220
292 Ile His Gly Arg Val Pro Tyr Ala Ile Ser Ile Ser Leu Ser Thr Ser
293 225 230 235 240
295 Arg Leu Lys Gln Val Gly Glu Lys Asn Leu Leu Lys Pro Leu Arg Asp
296 245 250 255
298 Thr Ala Glu Ala Ile Ser Arg Glu Leu Gly Phe Ser Val Arg Glu Gly
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304 <211> LENGTH: 332
305 <212> TYPE: PRT
306 <213> ORGANISM: YiaK-Ko
308 <400> SEQUENCE: 11
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313 Leu Asp Ala Cys Val Ala Arg Glu Thr Ala Asp Ala Cys Ala Glu Met

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